

A. BAKER

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11-21-00

P.2.

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/639,207

DATE: 11/06/2000
TIME: 11:21:18

Input Set : A:\Sequence Listing - 06618-686001.txt
Output Set: N:\CRF3\11062000\I639207.raw

4 <110> APPLICANT: Kazemi-Esfarjani, Parsa
5 Benzer, Seymour
7 <120> TITLE OF INVENTION: AN ANIMAL MODEL OF POLYGLUTAMINE
8 TOXICITY, METHODS OF USE, AND MODULATORS OF POLYGLUTAMINE
9 TOXICITY
11 <130> FILE REFERENCE: 06618-686001
13 <140> CURRENT APPLICATION NUMBER: US 09/639,207
14 <141> CURRENT FILING DATE: 2000-08-14
16 <150> PRIOR APPLICATION NUMBER: US 60/148,934
17 <151> PRIOR FILING DATE: 1999-08-12
19 <150> PRIOR APPLICATION NUMBER: US 60/148,933
20 <151> PRIOR FILING DATE: 1999-08-12
22 <150> PRIOR APPLICATION NUMBER: US 60/177,047
23 <151> PRIOR FILING DATE: 2000-01-18
25 <150> PRIOR APPLICATION NUMBER: US 60/205,720
26 <151> PRIOR FILING DATE: 2000-05-19
28 <160> NUMBER OF SEQ ID NOS: 69
30 <170> SOFTWARE: FastSEQ for Windows Version 4.0
32 <210> SEQ ID NO: 1
33 <211> LENGTH: 508
34 <212> TYPE: PRT
35 <213> ORGANISM: Human
37 <400> SEQUENCE: 1
38 Met Asp Asp Glu Val Ile Glu Ile Ser Asp Ser Glu Arg Glu Glu Thr
39 1 5 10 15
40 Ser Ser Asn Ser Glu Met Asp Val Glu Ile Thr Thr Glu Gln Pro Thr
41 20 25 30
42 Ile Asp Val Lys Ala Glu Gln Ile Val Pro Lys Asp Ala Ala Thr Ile
43 35 40 45
44 Ala Glu Glu Lys Lys Lys Leu Gly Asn Asp Gln Tyr Lys Ala Gln Asn
45 50 55 60
46 Tyr Gln Asn Ala Leu Lys Leu Tyr Thr Asp Ala Ile Ser Leu Cys Pro
47 65 70 75 80
48 Asp Ser Ala Ala Tyr Tyr Gly Asn Arg Ala Ala Cys Tyr Met Met Leu
49 85 90 95
50 Leu Asn Tyr Asn Ser Ala Leu Thr Asp Ala Arg His Ala Ile Arg Ile
51 100 105 110
52 Asp Pro Gly Phe Glu Lys Ala Tyr Val Arg Val Ala Lys Cys Cys Leu
53 115 120 125
54 Ala Leu Gly Asp Ile Ile Gly Thr Glu Gln Ala Val Lys Met Val Asn
55 130 135 140
56 Glu Leu Asn Ser Leu Ser Thr Ala Val Ala Ala Glu Gln Thr Ala Ala
57 145 150 155 160
58 Gln Lys Leu Arg Gln Leu Glu Ala Thr Ile Gln Ala Asn Tyr Asp Thr
59 165 170 175
60 Lys Ser Tyr Arg Asn Val Val Phe Tyr Leu Asp Ser Ala Leu Lys Leu
61 180 185 190

ENTERED

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62 Ala Pro Ala Cys Leu Lys Tyr Arg Leu Leu Lys Ala Glu Cys Leu Ala
63      195      200      205
64 Phe Leu Gly Arg Cys Asp Glu Ala Leu Asp Ile Ala Val Ser Val Met
65      210      215      220
66 Lys Leu Asp Thr Thr Ser Ala Asp Ala Ile Tyr Val Arg Gly Leu Cys
67 225      230      235      240
68 Leu Tyr Tyr Thr Asp Asn Leu Asp Lys Gly Ile Leu His Phe Glu Arg
69      245      250      255
70 Ala Leu Thr Leu Asp Pro Asp His Tyr Lys Ser Lys Gln Met Arg Ser
71      260      265      270
72 Lys Cys Lys Gln Leu Lys Glu Met Lys Glu Asn Gly Asn Met Leu Phe
73      275      280      285
74 Lys Ser Gly Arg Tyr Arg Glu Ala His Val Ile Tyr Thr Asp Ala Leu
75      290      295      300
76 Lys Ile Asp Glu His Asn Lys Asp Ile Asn Ser Lys Leu Leu Tyr Asn
77 305      310      315      320
78 Arg Ala Leu Val Asn Thr Arg Ile Gly Asn Leu Arg Glu Ala Val Ala
79      325      330      335
80 Asp Cys Asn Arg Val Leu Glu Leu Asn Ser Gln Tyr Leu Lys Ala Leu
81      340      345      350
82 Leu Leu Arg Ala Arg Cys Tyr Asn Asp Leu Glu Lys Phe Glu Glu Ser
83      355      360      365
84 Val Ala Asp Tyr Glu Thr Ala Leu Gln Leu Glu Lys Thr Pro Glu Ile
85      370      375      380
86 Lys Arg Met Leu Arg Glu Ala Lys Phe Ala Leu Lys Lys Ser Lys Arg
87 385      390      395      400
88 Lys Asp Tyr Tyr Lys Ile Leu Gly Ile Gly Arg Asn Ala Ser Asp Asp
89      405      410      415
90 Glu Ile Lys Lys Ala Tyr Arg Lys Lys Ala Leu Val His His Pro Asp
91      420      425      430
92 Arg His Ala Asn Ser Ser Ala Glu Glu Arg Lys Glu Glu Glu Leu Lys
93      435      440      445
94 Phe Lys Glu Val Gly Glu Ala Tyr Ala Ile Leu Ser Asp Ala His Lys
95      450      455      460
96 Lys Ser Arg Tyr Asp Ser Gly Gln Asp Ile Glu Glu Gln Glu Ala
97 465      470      475      480
98 Asp Phe Asp Pro Asn Gln Met Phe Arg Thr Phe Phe Gln Phe Asn Gly
99      485      490      495
100 Gly Gly Arg Asn Asn Ser Ser Phe Asn Phe Glu Phe
101      500      505
103 <210> SEQ ID NO: 2
104 <211> LENGTH: 2239
105 <212> TYPE: DNA
106 <213> ORGANISM: Drosophila
108 <400> SEQUENCE: 2
109 ggcacgagcc actacttcgc atggcacgct tttttccgtg tgcctcgggttc gttcgggcat      60
110 acaaaacaca aaattcaagt ttaaaaaacta aataggcaac taaaagggaa gccgcagcga      120
111 ataaagtgat ttgctgaaag agacgtaaga aagttaatcg catcgaaggc accagaaatc      180
112 ggggatttct aacacggcgc gcgtgcgacg tacatacata cgcaagcgca cacacacacg      240

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113 aacaattact tgcattgac gcaaaagcga aaaagcagtg gaataaaggg gaattgacaa 300
114 ataacaacgt ttgcaagca ctggaactctg gtcgctggtg ttcttlicatt ttgtaattgc 360
115 cagcagatga cgacgaagta attgaaatta ggcacagcga acgcgaagaa acctcactga 420
116 actccgaaat ggatgtgga ataacgacag aacagccaac cactgatgtc aaagcagagc 480
117 aaattgtjcc caaggacgag gcaaccattg ccgaggagaa gaagaaactg ggcaacgacc 540
118 aatacaaggc gcagaactat cagaatgcac tcaagctcta cagcgatgcc atatcgctgt 600
119 gtccggaactc ggcggcatatc tatggcaatc gggccgctcg ctacatgatg ctgctcaact 660
120 ataataagcgc cctgacacag gcccgacacg ccatacgcat cgatccgggc ttcgagaagg 720
121 cctacgtccg tgtggccaag tgcgtgtctg cctctggcga cattattggc accgaacagg 780
122 ccgtcaaaat ggtcaacgag ctgaattcgc ttacgacgac tgttgctgcc gaacagacgg 840
123 ggcgcgcaaaa gttgcgcgcaa ttggagcgca ccattcaggc aaactacgat acgaaatcct 900
124 atcgcaatgt ggtctctctat ttgatatgtg ccttgaaatt ggcgcgccgc tgtttgaaat 960
125 atcgtctact caaggctgag tgccttgcat ttttggggcg atgtgatgag gccttgagca 1020
126 ttgcggtcag tgaatgaaa ctggatacca catcgccgga tgcgatatac gtgagaggtc 1080
127 tgtgctgtga ctacacggac aacctggaca agggaattct tcatttcgag cgcgccctga 1140
128 cctcgacccc ggaacctac aaqtccaagc agatgcgcag caaatgcaag cagctcaagg 1200
129 agatgaagga gaacggcaat atgctattca agtcgggtcg gtatcgcgag gcacacgtta 1260
130 totacacgga cgccctgaag atcgatgaac acaacaagga tatcaattcg aaattgcttt 1320
131 acaatcgggc ttgtgtcaac acgcgtattg gcaatttcg agaggccgtg gccgattgca 1380
132 atcgagtgct ggagctgaat agtcagatc tgaaggctct gttgctgca ggcgcgtgct 1440
133 acaatgatct ggagaaagtc gaggagtcgg tggcgacta tgagacggcg ctgcagctgg 1500
134 agaagacgcc ggagattaa cgaatgctgc gcgaggccaa gtttgcgttg aagaagtcga 1560
135 agcgaagga ctactacaag atcctgggca ttggacgcaa tgcgtccgac gacgagatca 1620
136 agaagcgcta tcgcaaaaag gcgctgttac atcatccgga tcgacacgca aacagcagtg 1680
137 ccgaggagcg caaggaggag gagctcaagt tcaaggaggt ggcggaaggc tacgccatac 1740
138 tgtcggaatg tcacaagaag tcgcgtacg acagcgcca ggatacgag gacgaggagc 1800
139 aagccgactt cgatccgaat caaatgttcc gcacattctt ccaattcaac ggcggtggcc 1860
140 ggaataattc atcgttcaac tttgagttct aggatccaa cgagtgttgt tcaccaccac 1920
141 agagaagaag accatctcaa tccatactt tctgctcat ccgaaaccaa catacagcag 1980
142 cgcacaaatt ttgaactctt ttacatatct cttttccaaa aagcaagaaa ataccacatt 2040
143 ttgatlatgt taacgaatga atatatgcca agttatttga aaaaaatttc taaatcaaaa 2100
144 taatgcaact aaattttccag tgaagtcca cttttttaa tgttctttct tggatttttt 2160
145 ttctggcaac attataaat catgggagat ttgtgttaa taaacagaaa tatatatata 2220
146 aaaaaaaaaa aaaaaaaaaa 2239
148 <210> SEQ ID NO: 3
149 <211> LENGTH: 273
150 <212> TYPE: PRT
151 <213> ORGANISM: Drosophila
153 <400> SEQUENCE: 3
154 Met Ser Leu Phe Gly Ala Leu Met Gly Asp Phe Asp Asp Asp Leu Gly
155 1 5 10 15
156 Leu Met Asn Asn His Met Asn His Thr Met Asn Ala Met Asn Met Gln
157 20 25 30
158 Met Arg Ser Met Asn Arg Leu Met Asn Ser Phe Met Pro Asp Pro Phe
159 35 40 45
160 Met Gln Val Ser Pro Phe Asp Gln Gly Phe Gln Gln Asn Ala Leu Met
161 50 55 60
162 Glu Arg Pro Gln Met Pro Ala Met Pro Ala Met Gly Leu Phe Gly Met
163 65 70 75 80

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164 Pro Met Met Pro Asn Phe Asn Arg Leu Leu Asn Ala Asp Ile Gly Gly
165                               85                      90                      95
166 Asn Ser Gly Ala Ser Phe Cys Gln Ser Thr Val Met Thr Met Ser Ser
167                               100                     105                     110
168 Gly Pro Asp Gly Arg Pro Gln Ile Tyr Gln Ala Ser Thr Ser Thr Lys
169                               115                     120                     125
170 Thr Gly Pro Gly Gly Val Arg Glu Thr Arg Arg Thr Val Gln Asp Ser
171                               130                     135                     140
172 Arg Thr Gly Val Lys Lys Met Ala Ile Gly His His Ile Gly Glu Arg
173 145                               150                     155                     160
174 Ala His Ile Ile Glu Lys Glu Gln Asp Met Arg Ser Gly Gln Leu Glu
175                               165                     170                     175
176 Glu Arg Gln Glu Phe Ile Asn Leu Glu Glu Gly Glu Ala Glu Gln Phe
177                               180                     185                     190
178 Asp Arg Glu Phe Thr Ser Arg Ala Ser Arg Gly Ala Val Gln Ser Arg
179                               195                     200                     205
180 His His Ala Gly Gly Met Gln Ala Ile Met Pro Ala Arg Pro Ala Ala
181                               210                     215                     220
182 His Thr Ser Thr Leu Thr Ile Glu Pro Val Glu Asp Asp Asp Asp Asp
183 225                               230                     235                     240
184 Asp Asp Asp Cys Val Ile Gln Glu Gln Gln Pro Val Arg Ser Ser Ala
185                               245                     250                     255
186 Gly Arg His Tyr Ser Ser Ala Pro Thr Ala Pro Gln Asn Arg Tyr Asn
187                               260                     265                     270
188 Tyr

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191 <210> SEQ ID NO: 4

192 <211> LENGTH: 1753

193 <212> TYPE: DNA

194 <213> ORGANISM: Drosophila

196 <400> SEQUENCE: 4

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197 ggcacgagga aaatattcgt gaaaattctg catacggaaa gaagaaaatt cgagcaacag      60
198 aaagccaaca caatccacaa aaatgtcttt attcggagcg ttgatgggtg atttcgacga      120
199 cgatctcggc cttatgaaca accacatgaa ccacactatg aacgcgatga acatgcagat      180
200 gcgctcgatg aatcgctga tgaacagctt tatgcccgat cccttcacgc aggtctcgcc      240
201 ctttgaccag ggattccagc agaacgctct catggagcgt ccgcagatgc cggccatgcc      300
202 agccatgggc ctcttcggca tgcctatgat gccaaacttt aatcgctgt tgaacgctga      360
203 tattggtggc aattcaggcg catcctctct ccagagcacc gtgatgacca tgcctcggg      420
204 tcccgatggg cgtctcaga tctaccaggc cagcactagt accaaaacag gaccgggagg      480
205 cgttcgtgag acccgaggga cggtgaggga ctgcgcact ggggtgaaga agatggccat      540
206 tggtcacac atcggcgagc gggcacacat tattgagaaa gagcaggaca tgcgctcagg      600
207 acaactggag gagcgccagg agttcattaa tctggaggag ggagaagccg agcagtttga      660
208 cagggagttt acatcgcgcg ctatcgcgcg agcgggtgcag tcaagacatc atgctggttg      720
209 catgcaggcc atcatgcccg ccggtccagc ggcacacacc tcgacgttga ccattgagcc      780
210 agtggaggac gacgacgacg atgatgatga ctgtgtaac caggagcagc aaccggttcg      840
211 ctctcccgcg ggcgcgcat attccagtgc gccaacggca ccgcagaaca gatataatta      900
212 ctaaatctaa agtcaataca gtatatatta ctaactatcc gataaaaacg aaacagaatt      960
213 gcatactata aatttctgct aattacatc ccaactcggt tcaaacgaaa cgaatatcga      1020
214 atcgaaatca tagaatgcac agagcagcat acatccacat cctatgccg ccaatccgag      1080
215 gcgccaacaa cgtgccgtaa aacatttcca cacggaggac gaagcggccg gctcctacaa      1140

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216 ggcgggtcaag cgcggcaaga agaagtagta gaaacgtgat catctgtatg ccaacatctt 1200
217 ccgcacatcga cactcaaaaa cactaggaag caaagcgttg ggttctgttc catagcagga 1260
218 aaaccaattc aatatatttt taacaaacac aattctttac cagttctgtc ttatcctgcg 1320
219 tgagtcgacc agaatgcaac actaaaaaat gtacaacttc aagatgctat tgatgtgcac 1380
220 gcaggatata gaacaacttg cttaaattta cttaaaacaa atgtgactat tcaacgccga 1440
221 aatcattaca acacacactc tcagacctaa tcgaaaaatt caatgaaagt aatyggaatat 1500
222 atatgaaatc glaattataa qlttgaatta ttgattaat tctcaagttt ttagattttg 1560
223 ttagccacta agcttttaaat tatggatgcc agttagcgtg caaatgaaca caattgattt 1620
224 gaaggctccg aacgatagaa aacaacaatt accaatcccc caaatacatg taattcgtaa 1680
225 ggcctaagta aatgttaacg tgaatttaaat taaatggtaa ttacattata atagtaaaaa 1740
226 aaaaaaaaaa aaa 1753
228 <210> SEQ ID NO: 5
229 <211> LENGTH: 484
230 <212> TYPE: PRT
231 <213> ORGANISM: Human
233 <400> SEQUENCE: 5
234 Met Ala Ala Thr Glu Pro Glu Leu Leu Asp Asp Gln Glu Ala Lys Arg
235 1 5 10 15
236 Glu Ala Glu Thr Phe Lys Glu Gln Gly Asn Ala Tyr Tyr Ala Lys Lys
237 20 25 30
238 Asp Tyr Asn Glu Ala Tyr Asn Tyr Thr Lys Ala Ile Asp Met Cys
239 35 40 45
240 Pro Lys Asn Ala Ser Tyr Tyr Gly Asn Arg Ala Ala Thr Leu Met Met
241 50 55 60
242 Leu Gly Arg Phe Arg Glu Ala Leu Gly Asp Ala Gln Gln Ser Val Arg
243 65 70 75 80
244 Leu Asp Asp Ser Phe Val Arg Gly His Leu Arg Glu Gly Lys Cys His
245 85 90 95
246 Leu Ser Leu Gly Asn Ala Met Ala Ala Cys Arg Ser Phe Gln Arg Ala
247 100 105 110
248 Leu Glu Leu Asp His Lys Asn Ala Gln Ala Gln Gln Glu Phe Lys Asn
249 115 120 125
250 Ala Asn Ala Val Met Glu Tyr Glu Lys Ile Ala Glu Thr Asp Phe Glu
251 130 135 140
252 Lys Arg Asp Phe Arg Lys Val Val Phe Cys Met Asp Arg Ala Leu Glu
253 145 150 155 160
254 Phe Ala Pro Ala Cys His Arg Phe Lys Ile Leu Lys Ala Glu Cys Leu
255 165 170 175
256 Ala Met Leu Gly Arg Tyr Pro Glu Ala Gln Ser Val Ala Ser Asp Ile
257 180 185 190
258 Leu Arg Met Asp Ser Thr Asn Ala Asp Ala Leu Tyr Val Arg Gly Leu
259 195 200 205
260 Cys Leu Tyr Tyr Glu Asp Cys Ile Glu Lys Ala Val Gln Phe Phe Val
261 210 215 220
262 Gln Ala Leu Arg Met Ala Pro Asp His Glu Lys Ala Cys Ile Ala Cys
263 225 230 235 240
264 Arg Asn Ala Lys Ala Leu Lys Ala Lys Lys Glu Asp Gly Asn Lys Ala
265 245 250 255
266 Phe Lys Glu Gly Asn Tyr Lys Leu Ala Tyr Glu Leu Tyr Thr Glu Ala

```

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/639,207

DATE: 11/06/2000

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Input Set : A:\Sequence Listing - 06618-686001.txt

Output Set: N:\CRF3\11062000\I639207.raw

L:1278 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:16
L:1298 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:17
L:1338 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:19
L:1374 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:21
L:1429 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:24
L:1430 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:24
L:1455 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:25
L:1535 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:30
L:1540 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:30
L:1626 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:35
L:1649 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:36
L:1686 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:38
L:1761 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:42
L:1783 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:43
L:1840 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:46
L:1912 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:51
L:1917 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:51
L:1941 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:52
L:1961 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:53
L:2029 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:56
L:2140 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:62